



RAW SEQUENCE LISTING ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 10/803,575A
Source: IFWO
Date Processed by STIC: 6/17/04

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.

PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
- 2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY

FOR CRF SUBMISSION AND PATENTIN SOFTWARE QUESTIONS, PLEASE CONTACT MARK SPENCER, TELEPHONE: 571-272-2510; FAX: 571-273-0221

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER VERSION 4.2 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:

<http://www.uspto.gov/web/offices/pac/checker/chkrnote.htm>

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail.

Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom.

Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

1. EFS-Bio (<<http://www.uspto.gov/ebc/efs/downloads/documents.htm>> , EFS Submission User Manual - ePAVE)
2. U.S. Postal Service: Commissioner for Patents, P.O. Box 1450, Alexandria, VA 22313-1450
3. Hand Carry, Federal Express, United Parcel Service, or other delivery service (EFFECTIVE 06/05/04):
U.S. Patent and Trademark Office, 220 20th Street S., Customer Window, Mail Stop Sequence, Crystal Plaza Two, Lobby, Room 1B03, Arlington, VA 22202

Revised 05/17/04



IFWO

RAW SEQUENCE LISTING

DATE: 06/17/2004

PATENT APPLICATION: US/10/803,575A

TIME: 07:28:24

Input Set : A:\SYR-HDAC-5005-C1 replacement 01.ST25.txt

Output Set: N:\CRF4\06172004\J803575A.raw

3 <110> APPLICANT: Syrrx, Inc.
 5 <120> TITLE OF INVENTION: HISTONE DEACETYLASE INHIBITORS
 7 <130> FILE REFERENCE: SYR-HDAC-5005-C1
 9 <140> CURRENT APPLICATION NUMBER: US 10/803,575A
 10 <141> CURRENT FILING DATE: 2004-03-17
 12 <150> PRIOR APPLICATION NUMBER: US 60/455,437
 13 <151> PRIOR FILING DATE: 2003-03-17
 15 <150> PRIOR APPLICATION NUMBER: US 60/531,203
 16 <151> PRIOR FILING DATE: 2003-12-19
 18 <160> NUMBER OF SEQ ID NOS: 8
 20 <170> SOFTWARE: PatentIn version 3.2
 22 <210> SEQ ID NO: 1
 23 <211> LENGTH: 513
 24 <212> TYPE: PRT
 25 <213> ORGANISM: Artificial
 27 <220> FEATURE:
 28 <223> OTHER INFORMATION: Residues 1-482 of HDAC1 and a 6-histidine tag at the N-terminus

30 <400> SEQUENCE: 1
 32 Met Ser Tyr Tyr His His His His His His Asp Tyr Asp Ile Pro Thr
 33 1 5 10 15
 36 Thr Glu Asn Leu Tyr Phe Gln Gly Ala Met Glu Pro Gly Gly Ser Met
 37 20 25 30
 40 Ala Gln Thr Gln Gly Thr Arg Arg Lys Val Cys Tyr Tyr Tyr Asp Gly
 41 35 40 45
 44 Asp Val Gly Asn Tyr Tyr Tyr Gly Gln Gly His Pro Met Lys Pro His
 45 50 55 60
 48 Arg Ile Arg Met Thr His Asn Leu Leu Leu Asn Tyr Gly Leu Tyr Arg
 49 65 70 75 80
 52 Lys Met Glu Ile Tyr Arg Pro His Lys Ala Asn Ala Glu Glu Met Thr
 53 85 90 95
 56 Lys Tyr His Ser Asp Asp Tyr Ile Lys Phe Leu Arg Ser Ile Arg Pro
 57 100 105 110
 60 Asp Asn Met Ser Glu Tyr Ser Lys Gln Met Gln Arg Phe Asn Val Gly
 61 115 120 125
 64 Glu Asp Cys Pro Val Phe Asp Gly Leu Phe Glu Phe Cys Gln Leu Ser
 65 130 135 140
 68 Thr Gly Gly Ser Val Ala Ser Ala Val Lys Leu Asn Lys Gln Gln Thr
 69 145 150 155 160
 72 Asp Ile Ala Val Asn Trp Ala Gly Gly Leu His His Ala Lys Lys Ser
 73 165 170 175
 76 Glu Ala Ser Gly Phe Cys Tyr Val Asn Asp Ile Val Leu Ala Ile Leu
 77 180 185 190
 80 Glu Leu Leu Lys Tyr His Gln Arg Val Leu Tyr Ile Asp Ile Asp Ile

Does Not Comply
 Corrected Diskette Needed

(pg. 1-3, 5) 2

please explain the remaining residues

per
 the
 total
 length
 of the
 sequence.

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Input Set : A:\SYR-HDAC-5005-C1 replacement 01.ST25.txt

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```

81      195      200      205
84 His His Gly Asp Gly Val Glu Glu Ala Phe Tyr Thr Thr Asp Arg Val
85      210      215      220
88 Met Thr Val Ser Phe His Lys Tyr Gly Glu Tyr Phe Pro Gly Thr Gly
89 225      230      235      240
92 Asp Leu Arg Asp Ile Gly Ala Gly Lys Gly Lys Tyr Tyr Ala Val Asn
93      245      250      255
96 Tyr Pro Leu Arg Asp Gly Ile Asp Asp Glu Ser Tyr Glu Ala Ile Phe
97      260      265      270
100 Lys Pro Val Met Ser Lys Val Met Glu Met Phe Gln Pro Ser Ala Val
101      275      280      285
104 Val Leu Gln Cys Gly Ser Asp Ser Leu Ser Gly Asp Arg Leu Gly Cys
105      290      295      300
108 Phe Asn Leu Thr Ile Lys Gly His Ala Lys Cys Val Glu Phe Val Lys
109 305      310      315      320
112 Ser Phe Asn Leu Pro Met Leu Met Leu Gly Gly Gly Gly Tyr Thr Ile
113      325      330      335
116 Arg Asn Val Ala Arg Cys Trp Thr Tyr Glu Thr Ala Val Ala Leu Asp
117      340      345      350
120 Thr Glu Ile Pro Asn Glu Leu Pro Tyr Asn Asp Tyr Phe Glu Tyr Phe
121      355      360      365
124 Gly Pro Asp Phe Lys Leu His Ile Ser Pro Ser Asn Met Thr Asn Gln
125      370      375      380
128 Asn Thr Asn Glu Tyr Leu Glu Lys Ile Lys Gln Arg Leu Phe Glu Asn
129 385      390      395      400
132 Leu Arg Met Leu Pro His Ala Pro Gly Val Gln Met Gln Ala Ile Pro
133      405      410      415
136 Glu Asp Ala Ile Pro Glu Glu Ser Gly Asp Glu Asp Glu Asp Asp Pro
137      420      425      430
140 Asp Lys Arg Ile Ser Ile Cys Ser Ser Asp Lys Arg Ile Ala Cys Glu
141      435      440      445
144 Glu Glu Phe Ser Asp Ser Glu Glu Glu Gly Glu Gly Arg Lys Asn
145      450      455      460
148 Ser Ser Asn Phe Lys Lys Ala Lys Arg Val Lys Thr Glu Asp Glu Lys
149 465      470      475      480
152 Glu Lys Asp Pro Glu Glu Lys Lys Glu Val Thr Glu Glu Glu Lys Thr
153      485      490      495
156 Lys Glu Glu Lys Pro Glu Ala Lys Gly Val Lys Glu Glu Val Lys Leu
157      500      505      510

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160 Ala

164 <210> SEQ ID NO: 2

165 <211> LENGTH: 1542

166 <212> TYPE: DNA

167 <213> ORGANISM: Artificial

169 <220> FEATURE:

170 <223> OTHER INFORMATION: DNA sequence encoding residues 1-482 of HDAC1 and a 6-

histidine

171 tag at the N-terminus

173 <400> SEQUENCE: 2

174 atgtcgtact accatcacca tcaccatcac gattacgata tcccaacgac cgaaaacctg 60

Please explain
 the remaining
 residues per
 the
 total
 length
 of the
 sequence

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Input Set : A:\SYR-HDAC-5005-C1 replacement 01.ST25.txt

Output Set: N:\CRF4\06172004\J803575A.raw

```

176 tatttttcagg gcgccatgga acccggggga tccatggcgc agacgcaggg caccgcggagg 120
178 aaagtctgtt actactacga cggggatggt ggaaattact attatggaca aggccaccca 180
180 atgaagcctc accgaatccg catgactcat aatttgctgc tcaactatgg tctctaccga 240
182 aaaatggaaa tctatcgccc tcacaaagcc aatgctgagg agatgaccaa gtaccacagc 300
184 gatgactaca ttaaattctt gcgctccatc cgtccagata acatgtcggg gtacagcaag 360
186 cagatgcaga gattcaacgt tgggtaggac tgtccagtat tcgatggcct gtttgagttc 420
188 tgtcagttgt ctactggtgg ttctgtggca agtgctgtga aacttaataa gcagcagacg 480
190 gacatcgctg tgaattgggc tgggggcctg caccatgcaa agaagtcgga ggcacatcggc 540
192 ttctgttacg tcaatgatat cgtcttggcc atcctggaac tgctaaagta tcaccagagg 600
194 gtgctgtaca ttgacattga tattcaccat ggtgacggcg tggagaggcg cttctacacc 660
196 acggaccggg tcatgactgt gtcttttcat aagtatggag agtacttccc aggaactggg 720
198 gacctacggg atatcggggc tggcaaaggc aagtattatg ctgttaacta cccgctccga 780
200 gacgggattg atgacgagtc ctatgaggcc attttcaagc cggtcatgtc caaagtaatg 840
202 gagatgttcc agcctagtgc ggtggtctta cagtgtggct cagactccct atctggggat 900
204 cggttagggt gcttcaatct aactatcaaa ggacacgcca agtgtgtgga atttgtcaag 960
206 agctttaacc tgccatgct gatgctggga ggcggtggtt acaccattcg taacgttgcc 1020
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210 tacaatgact actttgaata ctttggacca gatttcaagc tccacatcag tccttccaat 1140
212 atgactaacc agaacacgaa tgagtacctg gagaagatca aacagcgact gtttgagaac 1200
214 cttagaatgc tgccgcacgc acctggggtc caaatgcagg cgattcctga ggacgccatc 1260
216 cctgaggaga gtggcgatga ggacgaagac gacctgaca agcgcattct gatctgctcc 1320
218 tctgacaaac gaattgcctg tgaggaagag ttctccgatt ctgaagagga gggagagggg 1380
220 ggcgcgaaga actcttccaa cttcaaaaaa gccaaagagag tcaaaaacaga ggatgaaaaa 1440
222 gagaaagacc cagaggagaa gaaagaagtc accgaaggag agaaaaccaa ggaggagaag 1500
224 ccagaagcca aaggggtcaa ggaggaggtc aagttggcct ga 1542

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227 <210> SEQ ID NO: 3

228 <211> LENGTH: 498

229 <212> TYPE: PRT

230 <213> ORGANISM: Artificial

232 <220> FEATURE:

233 <223> OTHER INFORMATION: Residues 1-488 of HDAC2 and a 6-histidine tag at the C-terminus

Same error

235 <400> SEQUENCE: 3

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237 Met Gly Ser Met Ala Tyr Ser Gln Gly Gly Gly Lys Lys Lys Val Cys
238 1 5 10 15
241 Tyr Tyr Tyr Asp Gly Asp Ile Gly Asn Tyr Tyr Tyr Gly Gln Gly His
242 20 25 30
245 Pro Met Lys Pro His Arg Ile Arg Met Thr His Asn Leu Leu Leu Asn
246 35 40 45
249 Tyr Gly Leu Tyr Arg Lys Met Glu Ile Tyr Arg Pro His Lys Ala Thr
250 50 55 60
253 Ala Glu Glu Met Thr Lys Tyr His Ser Asp Glu Tyr Ile Lys Phe Leu
254 65 70 75 80
257 Arg Ser Ile Arg Pro Asp Asn Met Ser Glu Tyr Ser Lys Gln Met Gln
258 85 90 95
261 Arg Phe Asn Val Gly Glu Asp Cys Pro Val Phe Asp Gly Leu Phe Glu
262 100 105 110
265 Phe Cys Gln Leu Ser Thr Gly Gly Ser Val Ala Gly Ala Val Lys Leu
266 115 120 125
269 Asn Arg Gln Gln Thr Asp Met Ala Val Asn Trp Ala Gly Gly Leu His

```

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Output Set: N:\CRF4\06172004\J803575A.raw

```

270      130      135      140
273 His Ala Lys Lys Ser Glu Ala Ser Gly Phe Cys Tyr Val Asn Asp Ile
274 145      150      155      160
277 Val Leu Ala Ile Leu Glu Leu Lys Tyr His Gln Arg Val Leu Tyr
278      165      170      175
281 Ile Asp Ile Asp Ile His His Gly Asp Gly Val Glu Glu Ala Phe Tyr
282      180      185      190
285 Thr Thr Asp Arg Val Met Thr Val Ser Phe His Lys Tyr Gly Glu Tyr
286      195      200      205
289 Phe Pro Gly Thr Gly Asp Leu Arg Asp Ile Gly Ala Gly Lys Gly Lys
290      210      215      220
293 Tyr Tyr Ala Val Asn Phe Pro Met Arg Asp Gly Ile Asp Asp Glu Ser
294 225      230      235      240
297 Tyr Gly Gln Ile Phe Lys Pro Ile Ile Ser Lys Val Met Glu Met Tyr
298      245      250      255
301 Gln Pro Ser Ala Val Val Leu Gln Cys Gly Ala Asp Ser Leu Ser Gly
302      260      265      270
305 Asp Arg Leu Gly Cys Phe Asn Leu Thr Val Lys Gly His Ala Lys Cys
306      275      280      285
309 Val Glu Val Val Lys Thr Phe Asn Leu Pro Leu Leu Met Leu Gly Gly
310      290      295      300
313 Gly Gly Tyr Thr Ile Arg Asn Val Ala Arg Cys Trp Thr Tyr Glu Thr
314 305      310      315      320
317 Ala Val Ala Leu Asp Cys Glu Ile Pro Asn Glu Leu Pro Tyr Asn Asp
318      325      330      335
321 Tyr Phe Glu Tyr Phe Gly Pro Asp Phe Lys Leu His Ile Ser Pro Ser
322      340      345      350
325 Asn Met Thr Asn Gln Asn Thr Pro Glu Tyr Met Glu Lys Ile Lys Gln
326      355      360      365
329 Arg Leu Phe Glu Asn Leu Arg Met Leu Pro His Ala Pro Gly Val Gln
330      370      375      380
333 Met Gln Ala Ile Pro Glu Asp Ala Val His Glu Asp Ser Gly Asp Glu
334 385      390      395      400
337 Asp Gly Glu Asp Pro Asp Lys Arg Ile Ser Ile Arg Ala Ser Asp Lys
338      405      410      415
341 Arg Ile Ala Cys Asp Glu Glu Phe Ser Asp Ser Glu Asp Glu Gly Glu
342      420      425      430
345 Gly Gly Arg Arg Asn Val Ala Asp His Lys Lys Gly Ala Lys Lys Ala
346      435      440      445
349 Arg Ile Glu Glu Asp Lys Lys Glu Thr Glu Asp Lys Lys Thr Asp Val
350      450      455      460
353 Lys Glu Glu Asp Lys Ser Lys Asp Asn Ser Gly Glu Lys Thr Asp Thr
354 465      470      475      480
357 Lys Gly Thr Lys Ser Glu Gln Leu Ser Asn Pro Gly His His His His
358      485      490      495
361 His His
365 <210> SEQ ID NO: 4
366 <211> LENGTH: 1497
367 <212> TYPE: DNA

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RAW SEQUENCE LISTING

DATE: 06/17/2004

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TIME: 07:28:24

Input Set : A:\SYR-HDAC-5005-C1 replacement 01.ST25.txt

Output Set: N:\CRF4\06172004\J803575A.raw

368 <213> ORGANISM: Artificial

370 <220> FEATURE:

371 <223> OTHER INFORMATION: DNA sequence encoding residues 1-488 of HDAC2 and a 6-histidine tag at the C-terminus

372 tag at the C-terminus

374 <400> SEQUENCE: 4

375 atgggatcca tggcgtagac tcaaggaggc ggcaaaaaaa aagtctgcta ctactacgac 60
 377 ggtgatattg gaaattatta ttatggacag ggtcatccca tgaagcctca tagaatccgc 120
 379 atgaccata acttgctgtt aaattatggc ttatacagaa aaatggaaat atataggccc 180
 381 cataaagcca ctgccgaaga aatgacaaaa tatcacagt atgagtatat caaatttcta 240
 383 cgggtcaataa gaccagataa catgtctgag tatagtaagc agatgcagag atttaagtgt 300
 385 ggagaagatt gtccagtgtt tgatggactc tttgagtttt gtcagctctc aactggcggt 360
 387 tcagttgctg gagctgtgaa gttaaaccga caacagactg atatggctgt taattgggct 420
 389 ggaggattac atcatgctaa gaaatcagaa gcacaggat tctgttacgt taatgatatt 480
 391 gtgcttgcca tccttgaatt actaaagtat catcagagag tcttatatat tgatatagat 540
 393 attcatcatg gtgatgggtg tgaagaagct ttttatacaa cagatcgtgt aatgacggta 600
 395 tcattccata aatatgggga atactttcct ggcacaggag acttgaggga tattggtgct 660
 397 ggaaaaggca aatactatgc tgtcaatttt ccaatgagag atggtataga tgatgagtca 720
 399 tatgggcaga tatttaagcc tattatctca aagggtgatg agatgtatca acctagtgtc 780
 401 gtggtattac agtgtggtgc agactcatta tctggtgata gactgggttg tttcaatcta 840
 403 acagtcaaag gtcattgctaa atgtgtagaa gttgtaaaaa cttttaactt accattactg 900
 405 atgcttgagg gaggtggcta cacaatccgt aatgttgctc gatgttgga atagagact 960
 407 gcagttgccc ttgatttgga gattcccaat gagttgccat ataatgatta ctttgagtat 1020
 409 tttggaccag acttcaaact gcatattagt ccttcaaaca tgacaaacca gaacactcca 1080
 411 gaatatatgg aaaagataaa acagcgtttg tttgaaaatt tgccgatgtt acctcatgca 1140
 413 cctggtgtcc agatgcaagc tattccagaa gatgctgttc atgaagacag tggagatgaa 1200
 415 gatggagaag atccagacaa gagaatttct attcgagcat cagacaagcg gatagcttgt 1260
 417 gatgaagaat tctcagattc tgaggatgaa ggagaaggag gtcgaagaaa tgtggctgat 1320
 419 cataagaaag gagcaaagaa agctagaatt gaagaagata agaaagaaac agaggacaaa 1380
 421 aaaacagacg ttaaggaaga agataaatcc aaggacaaca gtggtgaaaa aacagatacc 1440
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426 <210> SEQ ID NO: 5

427 <211> LENGTH: 782

428 <212> TYPE: PRT

429 <213> ORGANISM: Artificial

431 <220> FEATURE:

432 <223> OTHER INFORMATION: Residues 73-845 of HDAC6 and a 6-histidine tag at the C-terminus

434 <400> SEQUENCE: 5

436 Met Pro Gly Met Asp Leu Asn Leu Glu Ala Glu Ala Leu Ala Gly Thr
 437 1 5 10 15
 440 Gly Leu Val Leu Asp Glu Gln Leu Asn Glu Phe His Cys Leu Trp Asp
 441 20 25 30
 444 Asp Ser Phe Pro Glu Gly Pro Glu Arg Leu His Ala Ile Lys Glu Gln
 445 35 40 45
 448 Leu Ile Gln Glu Gly Leu Leu Asp Arg Cys Val Ser Phe Gln Ala Arg
 449 50 55 60
 452 Phe Ala Glu Lys Glu Glu Leu Met Leu Val His Ser Leu Glu Tyr Ile
 453 65 70 75 80
 456 Asp Leu Met Glu Thr Gln Tyr Met Asn Glu Gly Glu Leu Arg Val
 457 85 90 95

The type of errors shown exist throughout
 the Sequence Listing. Please check subsequent
 sequences for similar errors

RAW SEQUENCE LISTING ERROR SUMMARY DATE: 06/17/2004
PATENT APPLICATION: US/10/803,575A TIME: 07:28:25

Input Set : A:\SYR-HDAC-5005-C1 replacement 01.ST25.txt
Output Set: N:\CRF4\06172004\J803575A.raw

Invalid <213> Response:

Use of "Artificial" only as "<213> Organism" response is incomplete,
per 1.823(b) of New Sequence Rules. Valid response is Artificial Sequence.

Seq#:1,2,3,4,5,6,7,8

VERIFICATION SUMMARY

DATE: 06/17/2004

PATENT APPLICATION: US/10/803,575A

TIME: 07:28:25

Input Set : A:\SYR-HDAC-5005-C1 replacement 01.ST25.txt

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